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5-15-02
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RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/022,554A

DATE: 03/13/2002

TIME: 09:27:40

Input Set : A:\PTO.VSK.txt

Output Set: N:\CRF3\03132002\J022554A.raw

3 <110> APPLICANT: ETTENSON, David S.
 4 BURGESS, Wilson H.
 5 DROHAN, William N.
 6 MACIAG, Thomas
 8 <120> TITLE OF INVENTION: COMPOSITIONS, METHODS AND KITS RELATING TO THROMBIN
 DEGRADATION RESISTANT

9 FIBROBLAST GROWTH FACTOR-1

11 <130> FILE REFERENCE: 054474-5005

13 <140> CURRENT APPLICATION NUMBER: 10/022,554A

14 <141> CURRENT FILING DATE: 2001-12-17

16 <160> NUMBER OF SEQ ID NOS: 8

18 <170> SOFTWARE: PatentIn version 3.1

20 <210> SEQ ID NO: 1

21 <211> LENGTH: 638

22 <212> TYPE: DNA

23 <213> ORGANISM: Homo sapiens

25 <400> SEQUENCE: 1

26	gaattcggga	acgcgccaca	agcagcagct	gctgagccat	ggctgaagg	gaaatcacca	60
28	ccttcacagc	cctgaccgag	aagttaatac	tgccctccag	gaattacaag	aagcccaaac	120
30	tcctctactg	tagcaacggg	ggccacttcc	tgaggatcct	tccggatggc	acagtggatg	180
32	ggacaaggga	caggagcgac	cagcacattc	agctgcagct	cagtgcggaa	agcgtggggg	240
34	aggtgtatat	aaagagtacc	gagactggcc	agtacttggc	catggacacc	gacgggcttt	300
36	tatacggctc	acagacacca	aatgaggaat	gtttgttcct	ggaaaggctg	gaggagaacc	360
38	attacaacac	ctatatatcc	aagaagcatg	cagagaagaa	ttggtttggt	ggcctcaaga	420
40	agaatgggag	ctgcaaacgc	ggtcctcgga	ctcactatgg	ccagaaagca	atcttgtttc	480
42	tccccctgcc	agtctcttct	gattaaagag	atctgttctg	gtgttgacca	ctccagagaa	540
44	gtttcgaggg	gtcctcacct	ggttgacccc	aaaaatgttc	ccttgaccat	tggctgcgct	600
46	aacccccagc	ccacagagcc	tgaatttgta	agcaactt			638

49 <210> SEQ ID NO: 2

50 <211> LENGTH: 154

51 <212> TYPE: PRT

52 <213> ORGANISM: Homo sapiens

54 <400> SEQUENCE: 2

56	Ala	Glu	Gly	Glu	Ile	Thr	Thr	Phe	Thr	Ala	Leu	Thr	Glu	Lys	Phe	Asn
57	1			5						10					15	
60	Leu	Pro	Pro	Gly	Asn	Tyr	Lys	Lys	Pro	Lys	Leu	Leu	Tyr	Cys	Ser	Asn
61				20						25					30	
64	Gly	Gly	His	Phe	Leu	Arg	Ile	Leu	Pro	Asp	Gly	Thr	Val	Asp	Gly	Thr
65			35					40					45			
68	Arg	Asp	Arg	Ser	Asp	Gln	His	Ile	Gln	Leu	Gln	Leu	Ser	Ala	Glu	Ser
69		50					55					60				
72	Val	Gly	Glu	Val	Tyr	Ile	Lys	Ser	Thr	Glu	Thr	Gly	Gln	Tyr	Leu	Ala
73	65					70				75					80	
76	Met	Asp	Thr	Asp	Gly	Leu	Leu	Tyr	Gly	Ser	Gln	Thr	Pro	Asn	Glu	Glu

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77          85          90          95
80 Cys Leu Phe Leu Glu Arg Leu Glu Glu Asn His Tyr Asn Thr Tyr Ile
81          100          105          110
84 Ser Lys Lys His Ala Glu Lys Asn Trp Phe Val Gly Leu Lys Lys Asn
85          115          120          125
88 Gly Ser Cys Lys Arg Gly Pro Arg Thr His Tyr Gly Gln Lys Ala Ile
89          130          135          140
92 Leu Phe Leu Pro Leu Pro Val Ser Ser Asp
93 145          150
96 <210> SEQ ID NO: 3
97 <211> LENGTH: 638
98 <212> TYPE: DNA
99 <213> ORGANISM: Homo sapiens
101 <400> SEQUENCE: 3
102 gaattcggga acgcgccaca agcagcagct gctgagccat ggctgaaggg gaaatcacca      60
104 ccttcacagc cctgaccgag aagtttaatc tgcctccagg gaattacaag aagcccaaac      120
106 tcctctactg tagcaacggg ggccacttcc tgaggatcct tccggatggc acagtggatg      180
108 ggacaagggg caggagcgac cagcacattc agctgcagct cagtgcggaa agcgtggggg      240
110 aggtgtatat aaagagtacc gagactggcc agtacttggc catggacacc gacgggcttt      300
112 tatacggtc acagacacca aatgaggaat gtttgttcct ggaaaggctg gaggagaacc      360
114 attacaacac ctatatatcc aagaagcatg cagagaagaa ttggtttgtt ggcctcaaga      420
116 agaatgggag ctgcaaacgc ggtcctaaaa ctcactatgg ccagaaagca atcttgtttc      480
118 tccccctgcc agtctcttct gattaaagag atctgttctg gtgttgacca ctccagagaa      540
120 gtttcgaggg gtcctcacct ggttgacccc aaaaatgttc ccttgaccat tggctgcgct      600
122 aacccccagc ccacagagcc tgaatttgta agcaactt      638
125 <210> SEQ ID NO: 4
126 <211> LENGTH: 154
127 <212> TYPE: PRT
128 <213> ORGANISM: Homo sapiens
130 <400> SEQUENCE: 4
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133 1          5          10          15
136 Leu Pro Pro Gly Asn Tyr Lys Lys Pro Lys Leu Leu Tyr Cys Ser Asn
137          20          25          30
140 Gly Gly His Phe Leu Arg Ile Leu Pro Asp Gly Thr Val Asp Gly Thr
141          35          40          45
144 Arg Asp Arg Ser Asp Gln His Ile Gln Leu Gln Leu Ser Ala Glu Ser
145          50          55          60
148 Val Gly Glu Val Tyr Ile Lys Ser Thr Glu Thr Gly Gln Tyr Leu Ala
149 65          70          75          80
152 Met Asp Thr Asp Gly Leu Leu Tyr Gly Ser Gln Thr Pro Asn Glu Glu
153          85          90          95
156 Cys Leu Phe Leu Glu Arg Leu Glu Glu Asn His Tyr Asn Thr Tyr Ile
157          100          105          110
160 Ser Lys Lys His Ala Glu Lys Asn Trp Phe Val Gly Leu Lys Lys Asn
161          115          120          125
164 Gly Ser Cys Lys Arg Gly Pro Arg Thr His Tyr Gly Gln Lys Ala Ile
165          130          135          140
168 Leu Phe Leu Pro Leu Pro Val Ser Ser Asp

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172 <210> SEQ ID NO: 5
173 <211> LENGTH: 31
174 <212> TYPE: DNA
175 <213> ORGANISM: Artificial Sequence
177 <220> FEATURE:
178 <223> OTHER INFORMATION: PCR primer
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185 <211> LENGTH: 10
186 <212> TYPE: PRT
187 <213> ORGANISM: Artificial Sequence
189 <220> FEATURE:
190 <223> OTHER INFORMATION: Amino acid encoded by PCR primer
192 <400> SEQUENCE: 6
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195 1                               5                               10
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199 <211> LENGTH: 31
200 <212> TYPE: DNA
201 <213> ORGANISM: Artificial Sequence
203 <220> FEATURE:
204 <223> OTHER INFORMATION: PCR primer
206 <400> SEQUENCE: 7
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211 <211> LENGTH: 10
212 <212> TYPE: PRT
213 <213> ORGANISM: Artificial Sequence
215 <220> FEATURE:
216 <223> OTHER INFORMATION: PCR primer
218 <400> SEQUENCE: 8
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221 1                               5                               10

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RAW SEQUENCE LISTING ERROR SUMMARY
PATENT APPLICATION: US/10/022,554A

DATE: 03/13/2002
TIME: 09:27:41

Input Set : A:\PTO.VSK.txt
Output Set: N:\CRF3\03132002\J022554A.raw

Invalid Line Length:

The rules require that a line not exceed 72 characters in length. This includes spaces.

Seq#:1; Line(s) 8

VERIFICATION SUMMARY

DATE: 03/13/2002

PATENT APPLICATION: US/10/022,554A

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